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SEQUENCE LISTING

5 <110> Wolosker, Herman
Takashashi, Maasaki
Mothet, Jean-Pierre
Ferris, Christopher
Snyder, Solomon

10 <120> Mammalian Serine Protease

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<213> Mus musculus

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Sub
B1

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5 <212> DNA

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20 caggatcttt taagattcgt ggtgctctca atgccgtcag aagcttggtt cctgatgctt 240

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<400> 7

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<212> PRT

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35 40 45

Phe Gln Lys Thr Gly Ser Phe Lys Ile Arg Gly Ala Leu Asn Ala Ile

20 50 55 60

Arg Gly Leu Ile Pro Asp Thr Pro Glu Glu Lys Pro Lys Ala Val Val

65 70 75 80

Thr His Ser Ser Gly Asn His Gly Gln Ala Leu Thr Tyr Ala Ala Lys

85 90 95

25 Leu Glu Gly Ile Pro Ala Tyr Ile Val Val Pro Gln Thr Ala Pro Asn

100 105 110

Cys Lys Lys Leu Ala Ile Gln Ala Tyr Gly Ala Ser Ile Val Tyr Cys

115 120 125

Asp Pro Ser Asp Glu Ser Arg Glu Lys Val Thr Gln Arg Ile Met Gln

30 130 135 140

Glu Thr Glu Gly Ile Leu Val His Pro Asn Gln Glu Pro Ala Val Ile

145 150 155 160

Ala Gly Gln Gly Thr Ile Ala Leu Glu Val Leu Asn Gln Val Pro Leu

165 170 175

35 Val Asp Ala Leu Val Val Pro Val Gly Gly Gly Gly Met Val Ala Gly

180 185 190

Ile Ala Ile Thr Ile Lys Ala Leu Lys Pro Ser Val Lys Val Tyr Ala

195 200 205

Ala Glu Pro Ser Asn Ala Asp Asp Cys Tyr Gln Ser Lys Leu Lys Gly

40 210 215 220

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Glu Leu Thr Pro Asn Leu His Pro Pro Glu Thr Ile Ala Asp Gly Val
 225 230 235 240
 Lys Ser Ser Ile Gly Leu Asn Thr Trp Pro Ile Ile Arg Asp Leu Val
 245 250 255
 5 Asp Asp Val Phe Thr Val Thr Glu Asp Glu Ile Lys Tyr Ala Thr Gln
 260 265 270
 Leu Val Trp Gly Arg Met Lys Leu Leu Ile Glu Pro Thr Ala Gly Val
 275 280 285
 Ala Leu Ala Ala Val Leu Ser Gln His Phe Gln Thr Val Ser Pro Glu
 10 290 295 300
 Val Lys Asn Val Cys Ile Val Leu Ser Gly Gly Asn Val Asp Leu Thr
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 Ser Leu Asn Trp Val Gly Gln Ala Glu Arg Pro Ala Pro Tyr Gln Thr
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 15 Val Ser Val

<210> 9

<211> 1023

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<212> DNA

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 40 tctccataa cttgggtgaa gcaggctgaa agccagctt cttatcagtc tgtttctggt 1020

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taa

1023

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 <211> 340
 5 <212> PRT
 <213> Homo sapiens

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 20 25 30
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 35 40 45
 15 Phe Gln Lys Thr Gly Ser Phe Lys Ile Arg Gly Ala Leu Asn Ala Val
 50 55 60
 Arg Ser Leu Val Pro Asp Ala Leu Glu Arg Lys Pro Lys Ala Val Val
 65 70 75 80
 Thr His Ser Ser Gly Asn His Gly Gln Ala Leu Thr Tyr Ala Ala Lys
 20 85 90 95
 Leu Glu Gly Ile Pro Ala Tyr Ile Val Val Pro Gln Thr Ala Pro Asp
 100 105 110
 Cys Lys Lys Leu Ala Ile Gln Ala Tyr Gly Ala Ser Ile Val Tyr Cys
 115 120 125
 25 Glu Pro Ser Asp Glu Ser Arg Glu Asn Val Ala Lys Arg Val Thr Glu
 130 135 140
 Glu Thr Glu Gly Ile Met Val His Pro Asn Gln Glu Pro Ala Val Ile
 145 150 155 160
 Ala Gly Gln Gly Thr Ile Ala Leu Glu Val Leu Asn Gln Val Pro Leu
 30 165 170 175
 Val Asp Ala Leu Val Val Pro Val Gly Gly Gly Gly Met Leu Ala Gly
 180 185 190
 Ile Ala Ile Thr Val Lys Ala Leu Lys Pro Ser Val Lys Val Tyr Ala
 195 200 205
 35 Ala Glu Pro Ser Asn Ala Asp Asp Cys Tyr Gln Ser Lys Leu Lys Gly
 210 215 220
 Lys Leu Met Pro Asn Leu Tyr Pro Pro Glu Thr Ile Ala Asp Gly Val
 225 230 235 240
 Lys Ser Ser Ile Gly Leu Asn Thr Trp Pro Ile Ile Arg Asp Leu Val
 40 245 250 255

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Asp Asp Ile Phe Thr Val Thr Glu Asp Glu Ile Lys Cys Ala Thr Gln
260 265 270
Leu Val Trp Glu Arg Met Lys Leu Leu Ile Glu Pro Thr Ala Gly Val
275 280 285
5 Gly Val Ala Ala Val Leu Ser Gln His Phe Gln Thr Val Ser Pro Glu
290 295 300
Val Lys Asn Ile Cys Ile Val Leu Ser Gly Gly Asn Val Asp Leu Thr
305 310 315 320
Ser Ser Ile Thr Trp Val Lys Gln Ala Glu Arg Pro Ala Ser Tyr Gln
10 325 330 335
Ser Val Ser Val
340
